E CANCELLE MAY 0 4 2006 3024-119-SEQ.ST25 QUENCE LISTING Mermod, Nicolas <110> Puttini, Stefania Saugy, Damien Girod, Pierre Alain Bucher, Philipp Nguyen, Duc-Quang Calabrese, David <120> HIGH EFFICIENCY GENE TRANSFER AND EXPRESSION IN MAMMALIAN CELLS BY A MULTIPLE TRANSFECTION PROCEDURE OF MAR SEQUENCES <130> 3024-119 <140> 10/595,495 <141> 2006-04-24 <150> PCT/EP2004/011974 <151> 2004-10-22 <150> EP04002722.9 2004-02-06 <151> <150> US60/513,574 <151> 2003-10-24 <160> 241 <170> PatentIn version 3.3 <210> <211> 320 <212> DNA <213> Homo sapiens <220> <221> misc_binding <222> (1)..(320)<223> MAR of human chromosome 1, nt from 36686 to 37008 <220> <221> <222> misc_binding (1)..(320)<223> MAR of human chromosome 1, genomic contig; 36686 to 37008 <400> ttatattatg ttgttatata tattatatta tgttattaga ttatattatg ttgttatatt 60 120 taattatata ttacattata taatatataa taatattata taattata ttacattata 180 240 300 320 gtatataata ttatataata

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taatataaca atatattta tatattatat gttatatatt a Page 88	tatattata tataatataa 300

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misc_binding

<221>

<211> 427 <212> DNA <213> Homo sapiens

<223> MAR of chromosome 2 genomic contig; 60605573..60605999

<222> (1)..(427)

12237 MAR OF CITY OF MODELLE & GENERAL CONTROL OF MARCHES AND MARC	
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ttatatatta tatataatat atgttatata tattatatat tttatataat atatatat	120
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tttaatatta tctattatat attttatata atatatat	240
atatatattt tacataatat ataatatata atacgtatta tatataatat ataatacgta	300
ttttatataa tatataatac gtattatata taatacgtat tatatattat ataatata	360
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atatacatat ataaagtata tataatatat acatatataa agtatatata tcatatatac	180
atatataaag tatatatata atatatacat atatacatat ataaagtata tataacatat	240
atacatatat aaagtatata taacatatat acatatataa agtatatata taatatac	300
atatatacat atataaagta tatataacat atatacatat atacagtata tataacatat	360
•	

atacatatat acagtatata taacatatat acatatatac agtatatata acatatatac

atatatacag tatatataac atatatacat atatacatga agtatatata acatatatac

atatatacat gaagtatata taacatatat acatatata atgaagtata tataacatat atacatatat acatgaagta tatataacat atatacatat atacatata aaagtatata

taacatatac atatacat atataaagta taacatatac atatacat atataaagta

tatataatat ataacatata catatataaa gtatatataa tatataacat atacatatat

420

480

540

600

660

720

·	
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atataatata tatacatata tacatatata aagtatatat aatatatata catatataca	1020
tatataaagt atatataata tatatacata tatacatata taaagtatat ataatatata	1080
tacatatata catatataaa gtatatataa tatgtataca tatatacata tataaagtat	1140
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attatattat atataataaa tatatattat attatatata atatatattt atatataaat	180
atattatata taaatatata ttatatataa aatatttata tattat	240
tatatataaa tatttatata ttatatataa atatttatat attatatata aatatttata	300
tattatatat aaaatatatt atatatatta tatatattat	360
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ctatatatat ataactatat atataactat atatatat	300
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atatataact	atatatatat	aactatatat	aactatatat	atataactat	atatataact	420
atatatatat	aactatatat	ataactatat	atatataact	atatatataa	ctatatatat	480
ataactatat	atataactat	atatataact	atatatataa	ctatatatat	ataactatat	540
atataactat	atatatataa	ctatatatat	aactatatat	ataactatat	atataactat	600
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<210> 231 <211> 1486 <212> DNA

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ataatatata ttagatgtaa tataatatat taatatata tagatgtaat ataatatat	1260
aataatatat attagatata atataatata ttaataatat attagatata atataatata	1320
ttaataatat ataagatata atataatata ttaataatat ataagatata atataatata	1380
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	360
tttatatata taaatatata tattatatat tttaatatat aaaatatata tattat	420
tttaatatat aaaatatata tattatatat tttaatatat aaaatatata tattat	420

3024-119-SEQ.ST25 tttaatatat ataaaaatata tatattatat attttatata tattaaaatat atattttata	480
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ta	302
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attatattat acataatata ttatatataa tatattatat ataatattat	240
tataatatat ataataataa tattataata tataata	300
atatataatt ctatatataa tatatagaat tctatctatt tataatatat atagaattct	360
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gtatatatat gtgtgtatat acgtatatat acatatatac aatttttgta tatatacata	240
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tatataatat taatattata tattattata tattatatta tattaatatt atatatatat	180
aatatatata atatatataa tagtattata tataatatat ataatagtat tatatattat	240
atatatataa tactattata tatattatat ataatagtat tatatatatt atatatat	300
tactattata tataatatat actattatat aatatatat	360
tataatacta ttatatataa tatatataat actattatat ataatatata taatactatt	420
atatataata tatataatac tattatatat aatatatat	480
tataatacta ttatatataa tatatatatt atataaatt atattaatat ataata	540
catatataat aatagtatat ataatatata atatatat	600
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400 230	
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gtttatatat ataatgtgta tatataatgt ttatatata	240
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aatatatatt atataattat ataatatatt atgtattata taatatatat	300
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407	

3024-119-SEQ.ST25 ttatatatat aaacattata tatacaaatt atatatat	180
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attaatatat aatatataat agtatattat atacatatat aatatataca atatataata	420
†	421